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1	<u>CLAIMS</u>
2	What is claimed is:
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4	Claim 1. A biopolymer marker selected from the group
5	consisting of sequence ID (R)SNLDEDIIAEENIVSR(S),
6	(R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N)
7	or at least one analyte thereof useful in indicating at
8	least one particular disease state.
9	
10	Claim 2. The biopolymer marker of claim 1 wherein
11	said disease state is predictive of Alzheimers disease.
12	
13	Claim 3. A method for evidencing and categorizing at
14	least one disease state comprising:
15	obtaining a sample from a patient;
16	conducting mass spectrometric analysis on said
17	sample;
18	evidencing and categorizing at least one biopolymer
19	marker sequence or analyte thereof isolated from said
20	sample; and,
21	comparing said at least one isolated biopolymer
22	marker sequence or analyte thereof to the biopolymer
23	marker sequence as set forth in claim 1;
24	wherein correlation of said isolated himpolymer

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1	marker and said proporymer marker sequence as sec forch in
2	claim 1 evidences and categorizes said at least one
3	disease state.
4	
5	Claim 4. The method of claim 3, wherein said step
6	of evidencing and categorizing is particularly directed to
7	biopolymer markers or analytes thereof linked to at least
8	one risk of disease development of said patient.
9	
10	Claim 5. The method of claim 3, wherein said step
11	of evidencing and categorizing is particularly directed to
12	biopolymer markers or analytes thereof related to the
13	existence of a particular disease state.
14	
15	Claim 6. The method of claim 3, wherein the sample
16	is an unfractionated body fluid or a tissue sample.
17	
18	
19	Claim 7. The method of claim 3, wherein said sample
20	is at least one of the group consisting of blood, blood
21	products, urine, saliva, cerebrospinal fluid, and lymph.
22	
23	Claim 8. The method of claim 3, wherein said mass
24	spectrometric analysis is selected from the group

1 consisting of Surface Enhanced Laser Desorption Ionization 2 (SELDI) mass spectrometry (MS), Maldi Qq TOF, MS/MS, TOF-TOF, and ESI-Q-TOF or an ION-TRAP. 3 4 5 The method of claim 3, wherein said Claim 9. 6 patient is a human. 7 A diagnostic assay kit for determining 8 Claim 10. 9 the presence of the biopolymer marker or analyte thereof 10 of claim 1 comprising: 11 at least one biochemical material which is capable of 12 specifically binding with a biomolecule which includes at 13 least said biopolymer marker or analyte thereof, and 14 means for determining binding between said 15 biochemical material and said biomolecule; whereby at least one analysis to determine a presence 16 17 of a marker, analyte thereof, or a biochemical material 18 specific thereto, is carried out on a sample. 19 20 Claim 11. The diagnostic assay kit of claim 10, 21 wherein said biochemical material or biomolecule is 22 immobilized on a solid support. 23 24 Claim 12. The diagnostic assay kit of claim 10

A kit for diagnosing, determining risk-

assessment, and identifying therapeutic avenues related to

at least one labeled biochemical material.

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including:

Claim 18.

1	a disease state comprising.
2	at least one biochemical material which is capable of
3	specifically binding with a biomolecule which includes at
4	least one biopolymer marker selected from the group
5	consisting of sequence ID (R)SNLDEDIIAEENIVSR(S),
6	(R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N)
7	or at least one analyte thereof related to said disease
8	state; and
9	means for determining binding between said
10	biochemical material and said biomolecule;
11	whereby at least one analysis to determine a presence
12	of a marker, analyte thereof, or a biochemical material
13	specific thereto, is carried out on a sample.
14.	
15	Claim 19. The kit of claim 18, wherein said
16	biochemical material or biomolecule is immobilized on a
17	solid support.
18	
19	Claim 20. The kit of claim 18 including:
20	at least one labeled biochemical material.
21	
22	Claim 21. The kit of claim 18, wherein said
23	biochemical material is an antibody.
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1	Claim 22. The kit of claim 20, wherein said labeled
2	biochemical material is an antibody.
3	
4	Claim 23. The kit of claim 18, wherein the sample is
5	an unfractionated body fluid or a tissue sample.
6	
7	Claim 24. The kit of claim 18, wherein said sample
8	is at least one of the group consisting of blood, blood
9	products, urine, saliva, cerebrospinal fluid, and lymph.
10	
11	Claim 25. The kit of claim 18, wherein said
12.	biochemical material is at least one monoclonal antibody
13	specific therefore.
14	
15	Claim 26. The kit of claim 18, wherein said
16	diagnosing, determining risk assessment, and identifying
17	therapeutic avenues is carried out on a single sample.
18	
19	Claim 27. The kit of claim 18, wherein said
20	diagnosing, determining risk assessment, and identifying
21	therapeutic avenues is carried out on multiple samples
22	such that at least one analysis is carried out on a first
23	sample and at least another analysis is carried out on a
24	second sample.

1	Claim 28. The kit of claim 27, wherein said first
2	and second samples are obtained at different time periods
3	
4	Claim 29. Polyclonal antibodies produced against a
5	marker sequence ID selected from the group consisting of
6	sequence ID (R)SNLDEDIIAEENIVSR(S),
7	(R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N)
8	or at least one analyte thereof in at least one animal
9	host.
10	-
11	Claim 30. An antibody that specifically binds a
12	biopolymer including a marker selected from the group
13	consisting of sequence ID (R)SNLDEDIIAEENIVSR(S),
14	(R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N)
15	or at least one analyte thereof.
16	
17	Claim 31. The antibody of claim 30 that is a
18	monoclonal antibody.
19	
20	Claim 32. The antibody of claim 30 that is a
21	polyclonal antibody.
22	
23	Claim 33. A process for identifying therapeutic
24	avenues related to a disease state comprising:

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1 conducting an analysis as provided by the kit of claim 18; and 2 3 interacting with a biopolymer selected from the group 4 consisting of sequence ID (R) SNLDEDIIAEENIVSR(S), 5 (R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N) or at least one analyte thereof; 6 7 whereby therapeutic avenues are developed. 8 9 The process for identifying therapeutic 10 avenues related to a disease state in accordance with 11 claim 33, wherein said therapeutic avenues regulate the 12 presence or absence of the biopolymer selected from the 13 group consisting of sequence ID (R) SNLDEDIIAEENIVSR(S), 14 (R) EGVQKEDIPPADLSDQVPDTESETR(I), (K) VTIKPAPETEKRPQDAK(N) 15 or at least one analyte thereof. 16 17 Claim 35. The process for identifying therapeutic 18 avenues related to a disease state in accordance with 19 claim 33, wherein said therapeutic avenues developed 20 include at least one avenue selected from a group 21 consisting of 1)utilization and recognition of said 22 biopolymer markers, variants or moieties thereof as direct 23 therapeutic modalities, either alone or in conjunction with an effective amount of a pharmaceutically effective 24

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carrier; 2) validation of therapeutic modalities or disease 2 preventative agents as a function of biopolymer marker 3 presence or concentration; 3) treatment or prevention of a 4 disease state by formation of disease intervention 5 modalities; 4) use of biopolymer markers or moieties 6 thereof as a means of elucidating therapeutically viable 7 agents, 5) instigation of a therapeutic immunological response; and 6) synthesis of molecular structures related 8 9 to said biopolymer markers, moieties or variants thereof 10 which are constructed and arranged to therapeutically 11 intervene in said disease state. 12 13 The process for identifying therapeutic Claim 36. 14 avenues related to a disease state in accordance with 15 claim 35, wherein said treatment or prevention of a 16 disease state by formation of disease intervention 17 modalities is the formation of biopolymer/ligand 18 conjugates which intervene at receptor sites to prevent, 19 delay or reverse a disease process. 20 21 Claim 37. The process for identifying therapeutic 22 avenues related to a disease state in accordance with 23 claim 35, wherein said means of elucidating 24 therapeutically viable agents includes use of a

1	bacteriophage peptide display library or a bacteriophage
2	antibody library.
3	
4	Claim 38. A process for regulating a disease state
5	by controlling the presence or absence of a biopolymer
6	selected from the group consisting of sequence ID
7	(R) SNLDEDIIAEENIVSR(S), (R) EGVQKEDIPPADLSDQVPDTESETR(I),
8	(K) $VTIKPAPETEKRPQDAK(N)$ or at least one analyte thereof.
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